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R045FR11 APP
LISTE DE SEQUENCES

<110> CYTOMICS SYSTEMS

<120> Procédé de criblage in vitro d'agents modulant
l'ubiquitination de la protéine I-Kappa-B-Alpha et
moyens destinés à la mise en oeuvre dudit procédé

<130> CYTOMICS

<140>
<141>

<160> 25

<170> PatentIn Ver. 2.1

<210> 1
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<213> Séquence artificielle

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<223> Description de la séquence
artificielle: GFP-NLS-IkBa

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<223> Description de la séquence
artificielle:GFP-NLS-IkBa

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Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile Cys
35 40 45

Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr Phe
50 55 60

Gly Tyr Gly Val Gln Cys Phe Ala Arg Tyr Pro Asp His Met Lys Gln
65 70 75 80

His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu Arg
85 90 95

Thr Ile Phe Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu Val
100 105 110

Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile
115 120 125

Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn
130 135 140

Tyr Asn Ser His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly
145 150 155 160

Ile Lys Val Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Ser Val
165 170 175

Gln Leu Ala Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro
180 185 190

Val Leu Leu Pro Asp Asn His Tyr Leu Ser Thr Gln Ser Ala Leu Ser
195 200 205

Lys Asp Pro Asn Glu Lys Arg Asp His Met Val Leu Leu Glu Phe Val
210 215 220

Thr Ala Ala Gly Ile Thr His Gly Met Asp Glu Leu Tyr Lys Leu Gln
225 230 235 240

Ser Pro Pro Pro Lys Lys Arg Lys Val Glu Leu Gly Gly Ser Met
245 250 255

Phe Gln Ala Ala Glu Arg Pro Gln Glu Trp Ala Met Glu Gly Pro Arg
260 265 270

Asp Gly Leu Lys Lys Glu Arg Leu Leu Asp Asp Arg His Asp Ser Gly
275 280 285

Leu Asp Ser Met Lys Asp Glu Glu Tyr Glu Gln Met Val Lys Glu Leu
290 295 300

Gln Glu Ile Arg Leu Glu Pro Gln Glu Val Pro Arg Gly Ser Glu Pro

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Trp Lys Gln Gln Leu Thr Glu Asp Gly Asp Ser Phe Leu His Leu Ala			
325	330	335	
Ile Ile His Glu Glu Lys Ala Leu Thr Met Glu Val Ile Arg Gln Val			
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Lys Gly Asp Leu Ala Phe Leu Asn Phe Gln Asn Asn Leu Gln Gln Thr			
355	360	365	
Pro Leu His Leu Ala Val Ile Thr Asn Gln Pro Glu Ile Ala Glu Ala			
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artificielle:GFP-NLS-bTRCP

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Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile Cys
35 40 45

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Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr Phe
 50 55 60
 Gly Tyr Gly Val Gln Cys Phe Ala Arg Tyr Pro Asp His Met Lys Gln
 65 70 75 80
 His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu Arg
 85 90 95
 Thr Ile Phe Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu Val
 100 105 110
 Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile
 115 120 125
 Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn
 130 135 140
 Tyr Asn Ser His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly
 145 150 155 160
 Ile Lys Val Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Ser Val
 165 170 175
 Gln Leu Ala Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro
 180 185 190
 Val Leu Leu Pro Asp Asn His Tyr Leu Ser Thr Gln Ser Ala Leu Ser
 195 200 205
 Lys Asp Pro Asn Glu Lys Arg Asp His Met Val Leu Leu Glu Phe Val
 210 215 220
 Thr Ala Ala Gly Ile Thr His Gly Met Asp Glu Leu Tyr Lys Leu Gln
 225 230 235 240
 Ser Pro Pro Pro Lys Lys Arg Lys Val Glu Leu Gly Gly Ser Met
 245 250 255
 Asp Pro Ala Glu Ala Val Leu Gln Glu Lys Ala Leu Lys Phe Met Cys
 260 265 270
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 275 280 285
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 Ala Phe Gln Asn Ser Ser Glu Arg Glu Asp Cys Asn Asn Gly Glu Pro
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 Pro Arg Lys Ile Ile Pro Glu Lys Asn Ser Leu Arg Gln Thr Tyr Asn
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 Ser Cys Ala Arg Leu Cys Leu Asn Gln Glu Thr Val Cys Leu Ala Ser
 340 345 350
 Thr Ala Met Lys Thr Glu Asn Cys Val Ala Lys Thr Lys Leu Ala Asn
 355 360 365
 Gly Thr Ser Ser Met Ile Val Pro Lys Gln Arg Lys Leu Ser Ala Ser
 370 375 380

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Tyr Glu Lys Glu Lys Glu Leu Cys Val Lys Tyr Phe Glu Gln Trp Ser
385 390 395 400

Glu Ser Asp Gln Val Glu Phe Val Glu His Leu Ile Ser Gln Met Cys
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His Tyr Gln His Gly His Ile Asn Ser Tyr Leu Lys Pro Met Leu Gln
420 425 430

Arg Asp Phe Ile Thr Ala Leu Pro Ala Arg Gly Leu Asp His Ile Ala
435 440 445

Glu Asn Ile Leu Ser Tyr Leu Asp Ala Lys Ser Leu Cys Ala Ala Glu
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565 570 575

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660 665 670

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Phe Asp Asp Lys Tyr Ile Val Ser Ala Ser Gly Asp Arg Thr Ile Lys
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Val Trp Asn Thr Ser Thr Cys Glu Phe Val Arg Thr Leu Asn Gly His
705 710 715 720

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Phe Asp Asn Lys Arg Ile Val Ser Gly Ala Tyr Asp Gly Lys Ile Lys
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785 790 795 800
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Gln Phe Asp Glu Phe Gln Ile Val Ser Ser Ser His Asp Asp Thr Ile
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850 855 860

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<212> PRT

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<220>
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<210> 19
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Leu Asp Thr

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